RAW SEQUENCE LISTING PATENT APPLICATION US/08/366,083A

DATE: 09/09/98 TIME: 10:15:36

INPUT SET: S28453.raw

This Raw Listing contains the General Information Section and up to the first 5 pages.

1	SEQUENCE LISTING ENTED
2	
3	(1) General Information:
4	
5	(i) APPLICANT: Pomerantz, Joel L.
6	Sharp, Phillip A.
7	Pabo, Carl O.
8	A A A MITTER OF INVENTION OUTVERING MENNIGHTEM ON TRACKORS
•9 10	(ii) TITLE OF INVENTION: CHIMERIC TRANSCRIPTION FACTORS
11	(iii) NUMBER OF SEQUENCES: 32
12	(III) NOMBER OF SEQUENCES: 32
13	(iv) CORRESPONDENCE ADDRESS:
14	(A) ADDRESSEE: FOLEY, HOAG & ELIOT LLP
15	(B) STREET: One Post Office Square
16	(C) CITY: Boston
17	(D) STATE: MA
18	(E) COUNTRY: USA
19	(F) ZIP: 02109-2170
20	
21	(V) COMPUTER READABLE FORM:
22	(A) MEDIUM TYPE: Floppy disk
23	(B) COMPUTER: IBM PC compatible
24	(C) OPERATING SYSTEM: PC-DOS/MS-DOS
25	(D) SOFTWARE: PatentIn Release #1.0, Version #1.30
26	() GUDDENM ADDITGAMION DAMA.
27 28	<pre>(vi) CURRENT APPLICATION DATA: (A) APPLICATION NUMBER: US 08/366,083</pre>
29	(B) FILING DATE: 29-DEC-1994
30	(C) CLASSIFICATION:
31	(C) CHADDITICATION.
32	(viii) ATTORNEY/AGENT INFORMATION:
33	(A) NAME: Vincent, Matthew P.
34	(B) REGISTRATION NUMBER: 36,709
35	(C) REFERENCE/DOCKET NUMBER: APV-022.01
36	
37	(ix) TELECOMMUNICATION INFORMATION:
38	(A) TELEPHONE: 617-832-1000
39	(B) TELEFAX: 617-832-7000
40	
41	AND THE PROPERTY HOP ONE TO WO I
42	(2) INFORMATION FOR SEQ ID NO:1:
43 44	(i) SPOURNOR OUADACHEDISTICS.
44	(i) SEQUENCE CHARACTERISTICS:(A) LENGTH: 26 base pairs
46	(A) LENGTH: 26 base pairs (B) TYPE: nucleic acid
40	(b) TIPE. NUCLEIC ACTO

RAW SEQUENCE LISTING PATENT APPLICATION US/08/366,083A

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		101 0 1 3E1. 320733.1um
47	(C) STRANDEDNESS: single	
48	(D) TOPOLOGY: linear	
49		
50	(ii) MOLECULE TYPE: DNA (genomic)	
51	(3)	
52		
53		
54	A CONTRACT PROGRAMMENT OF THE VALUE OF THE V	
55	(xi) SEQUENCE DESCRIPTION: SEQ ID NO:1:	
56		
57	GTTTGGCACC TGACTAATTT AAGGAG	26
58		
59	(2) INFORMATION FOR SEQ ID NO:2:	
60	-	
61	(i) SEQUENCE CHARACTERISTICS:	
62	(A) LENGTH: 25 base pairs	
	· · · · · · · · · · · · · · · · · · ·	
63	(B) TYPE: nucleic acid	
64	(C) STRANDEDNESS: single	
65	(D) TOPOLOGY: linear	
66		
67	(ii) MOLECULE TYPE: DNA (genomic)	
68		
69		
70		
71		
72	(xi) SEQUENCE DESCRIPTION: SEQ ID NO:2:	
73	(XI) SEQUENCE DESCRIPTION: SEQ ID NO.2:	
	CCCMTTANTA ACCCACCOTA CCCCC	25
74	GCGTTAATTA AGGGAGGTAA GGCCC	25
75		
76	(2) INFORMATION FOR SEQ ID NO:3:	
77		
78	(i) SEQUENCE CHARACTERISTICS:	
79	(A) LENGTH: 25 base pairs	
80	(B) TYPE: nucleic acid	
81	(C) STRANDEDNESS: single	
82	(D) TOPOLOGY: linear	
83	(b) Torobosi: Illieur	
84	(ii) MOLECULE TYPE: DNA (conomic)	
	(ii) MOLECULE TYPE: DNA (genomic)	
85		
86		
87		
88		
89	(xi) SEQUENCE DESCRIPTION: SEQ ID NO:3:	
90		
91	CTCGGCCGTT AATGAGGGGT GTTCG	25
92		
93	(2) INFORMATION FOR SEQ ID NO:4:	
94	(a) The ordination for phy in no.11	
	(i) GEOLIENGE GUADAGMEDIGMIGG	
95	(i) SEQUENCE CHARACTERISTICS:	
96	(A) LENGTH: 25 base pairs	
97	(B) TYPE: nucleic acid	
98	(C) STRANDEDNESS: single	
99	(D) TOPOLOGY: linear	
	• •	

RAW SEQUENCE LISTING PATENT APPLICATION US/08/366,083A

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INPUT SET: S28453, raw (ii) MOLECULE TYPE: DNA (genomic) (xi) SEQUENCE DESCRIPTION: SEO ID NO:4: TAATTATGGG CGGGATCGAA TAGCC (2) INFORMATION FOR SEQ ID NO:5: (i) SEQUENCE CHARACTERISTICS: (A) LENGTH: 26 base pairs (B) TYPE: nucleic acid (C) STRANDEDNESS: single (D) TOPOLOGY: linear (ii) MOLECULE TYPE: DNA (genomic) (xi) SEQUENCE DESCRIPTION: SEQ ID NO:5: GGCAATAATC AATCCTTTAA TTATGG (2) INFORMATION FOR SEQ ID NO:6: (i) SEQUENCE CHARACTERISTICS: (A) LENGTH: 26 base pairs (B) TYPE: nucleic acid (C) STRANDEDNESS: single (D) TOPOLOGY: linear (ii) MOLECULE TYPE: DNA (genomic) (xi) SEQUENCE DESCRIPTION: SEQ ID NO:6: GGCCGTACCT CATGAAATTA GGGGCG (2) INFORMATION FOR SEQ ID NO:7: (i) SEQUENCE CHARACTERISTICS: (A) LENGTH: 25 base pairs (B) TYPE: nucleic acid (C) STRANDEDNESS: single (D) TOPOLOGY: linear

(ii) MOLECULE TYPE: DNA (genomic)

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153		
154		
155		
156		
157	(xi) SEQUENCE DESCRIPTION: SEQ ID NO:7:	
158		
159	GTTAATTATG GGGTAATAAT GGTGC	25
160		
161	(2) INFORMATION FOR SEQ ID NO:8:	
162		
163	(i) SEQUENCE CHARACTERISTICS:	
164	(A) LENGTH: 25 base pairs	
165	(B) TYPE: nucleic acid	
166	(C) STRANDEDNESS: single	
167	(D) TOPOLOGY: linear	
168		
169	(ii) MOLECULE TYPE: DNA (genomic)	
170		
171		
172		
173		
174	(xi) SEQUENCE DESCRIPTION: SEQ ID NO:8:	
175	(AI) Digolated biboart flow big in nove.	
176	GTCGGGCTCT GTTAATTATG GGTGG	25
177	Greaderer Grigarian Garde	
178	(2) INFORMATION FOR SEQ ID NO:9:	
179	(2) INFORMATION FOR SEQ 15 No. 3.	
180	(i) SEQUENCE CHARACTERISTICS:	
181	(A) LENGTH: 25 base pairs	
182	(A) LENGTH: 25 base parts (B) TYPE: nucleic acid	
183	(C) STRANDEDNESS: single	
184	(D) TOPOLOGY: linear	
185	(ii) NOI EGUI E EVEE DNA (Gonomia)	
186	(ii) MOLECULE TYPE: DNA (genomic)	
187		
188		
189		
190		
191	(xi) SEQUENCE DESCRIPTION: SEQ ID NO:9:	
192		0.5
193	GGATAATTAC GGGTGGCATT TAGGC	25
194		
195	(2) INFORMATION FOR SEQ ID NO:10:	
196		
197	(i) SEQUENCE CHARACTERISTICS:	
198	(A) LENGTH: 25 base pairs	
199	(B) TYPE: nucleic acid	
200	(C) STRANDEDNESS: single	
201	(D) TOPOLOGY: linear	
202		
203	(ii) MOLECULE TYPE: DNA (genomic)	
204		
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		INFOT 3E1. 320433.74W
206		
207		
208	(xi) SEQUENCE DESCRIPTION: SEQ ID NO:10:	
209		
210	GATAAATAGG GGCGTCCCAT CCCGT	25
211		
212	(2) INFORMATION FOR SEQ ID NO:11:	
213		
214	(i) SEQUENCE CHARACTERISTICS:	
215	(A) LENGTH: 24 base pairs	
216	(B) TYPE: nucleic acid	
217	(C) STRANDEDNESS: single	
218	(D) TOPOLOGY: linear	
219		
220	(ii) MOLECULE TYPE: DNA (genomic)	
221		
222		
223		
224		
225	(xi) SEQUENCE DESCRIPTION: SEQ ID NO:11:	
226		
227	TAAATTAGGG CTTTAATTAC GGTC	24
228		
229	(2) INFORMATION FOR SEQ ID NO:12:	
230		
231	(i) SEQUENCE CHARACTERISTICS:	
232	(A) LENGTH: 25 base pairs	
233	(B) TYPE: nucleic acid	
234	(C) STRANDEDNESS: single	
235	(D) TOPOLOGY: linear	
236		
237	(ii) MOLECULE TYPE: DNA (genomic)	
238		
239		
240		
241		
242	(xi) SEQUENCE DESCRIPTION: SEQ ID NO:12:	
243	manmanan ammanan massa	25
244	TCATTAGAGT GTTAATGAGA TGCGC	25
245	(2) INFORMATION FOR CEO ID NO.12.	
246	(2) INFORMATION FOR SEQ ID NO:13:	
247	(i) CEOUENCE CUNDACTEDICATOR.	
248	(i) SEQUENCE CHARACTERISTICS:	
249	(A) LENGTH: 26 base pairs(B) TYPE: nucleic acid	
250	, ,	
251 252	<pre>(C) STRANDEDNESS: single (D) TOPOLOGY: linear</pre>	
252 253	(D) TOPOLOGI: IIMedi	
253 254	(ii) MOLECULE TYPE: DNA (genomic)	
254 255	(II) MODECODE TIPE. DNA (Genomic)	
255 256		
257		
257 258		
230		

SEQUENCE VERIFICATION REPORT PATENT APPLICATION *US/08/366,083A*

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